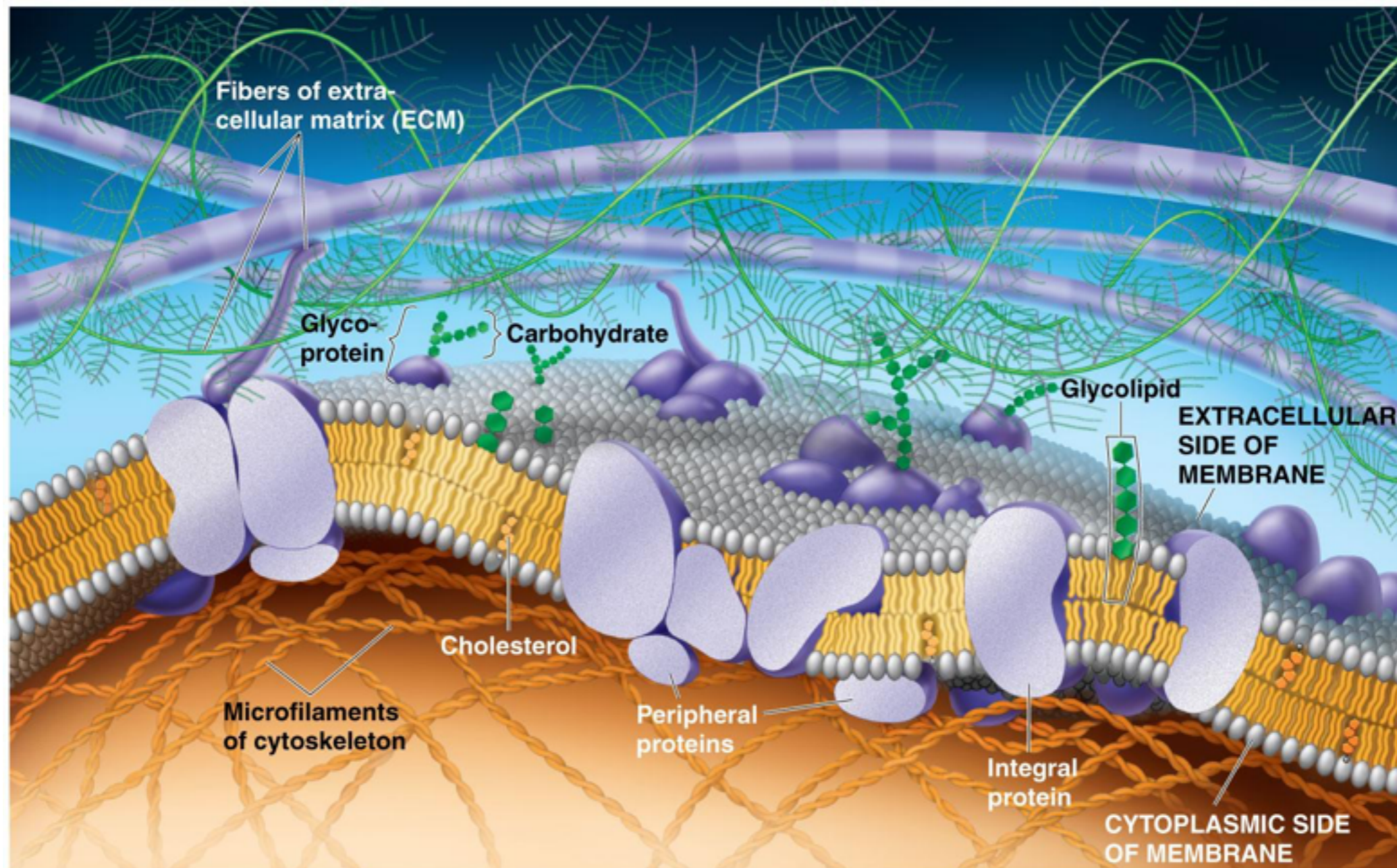


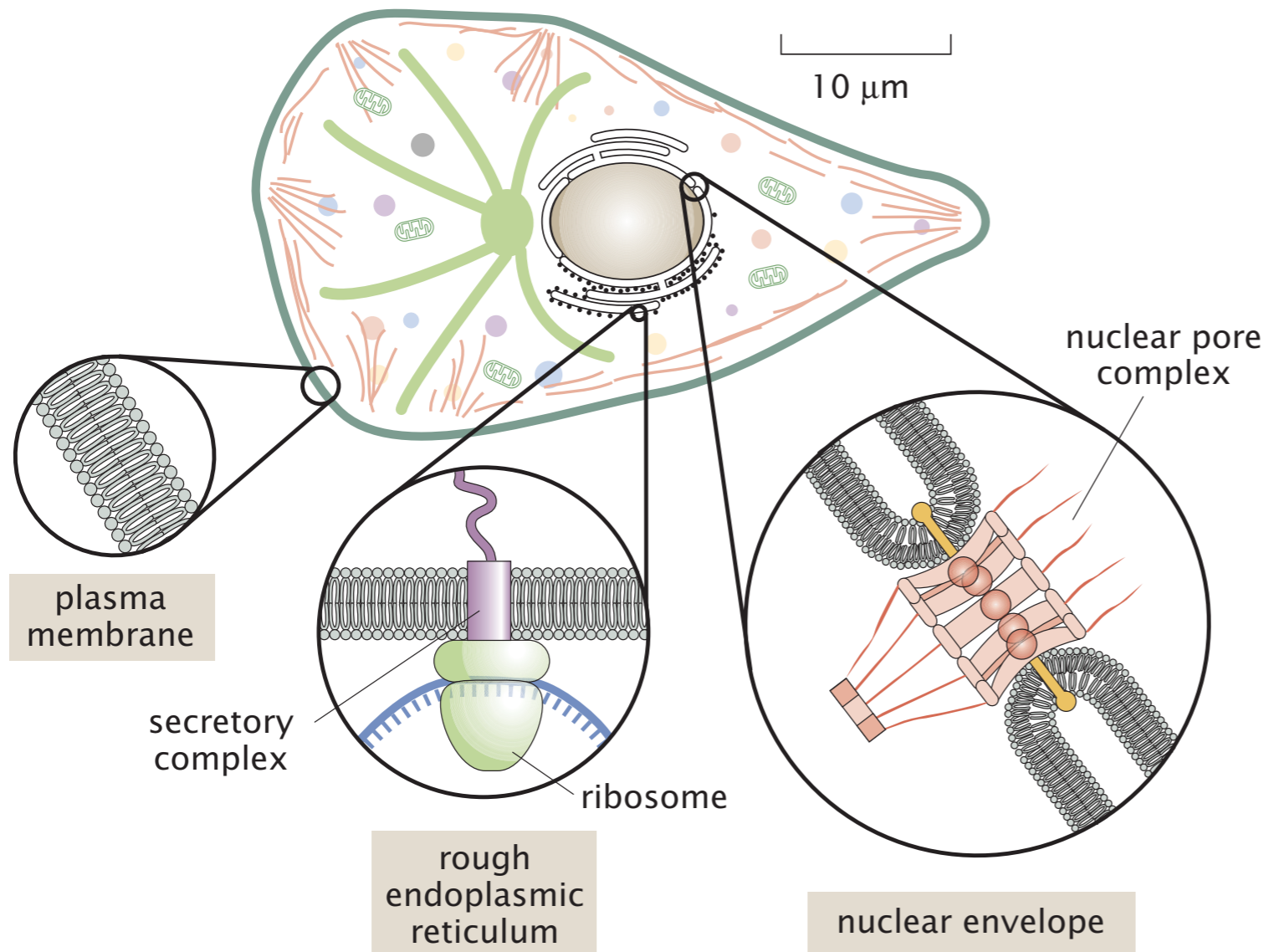
MAE 545: Lecture 14 (11/10)

Mechanics of cell membranes

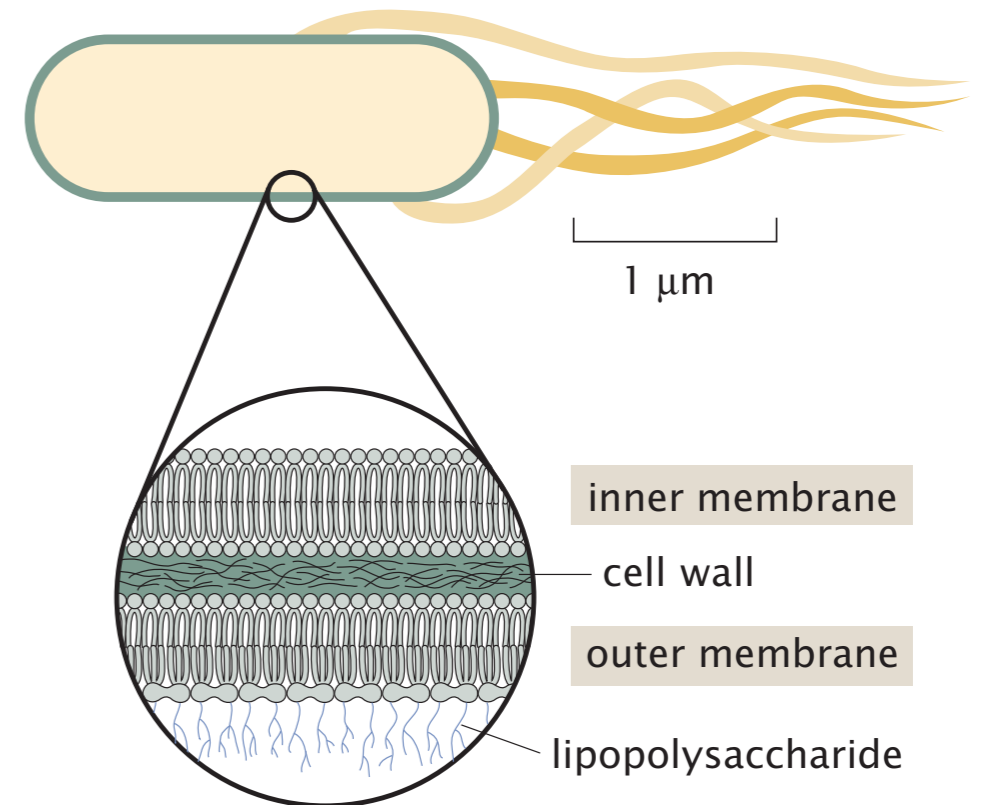


Cell membranes

Eukaryotic cells

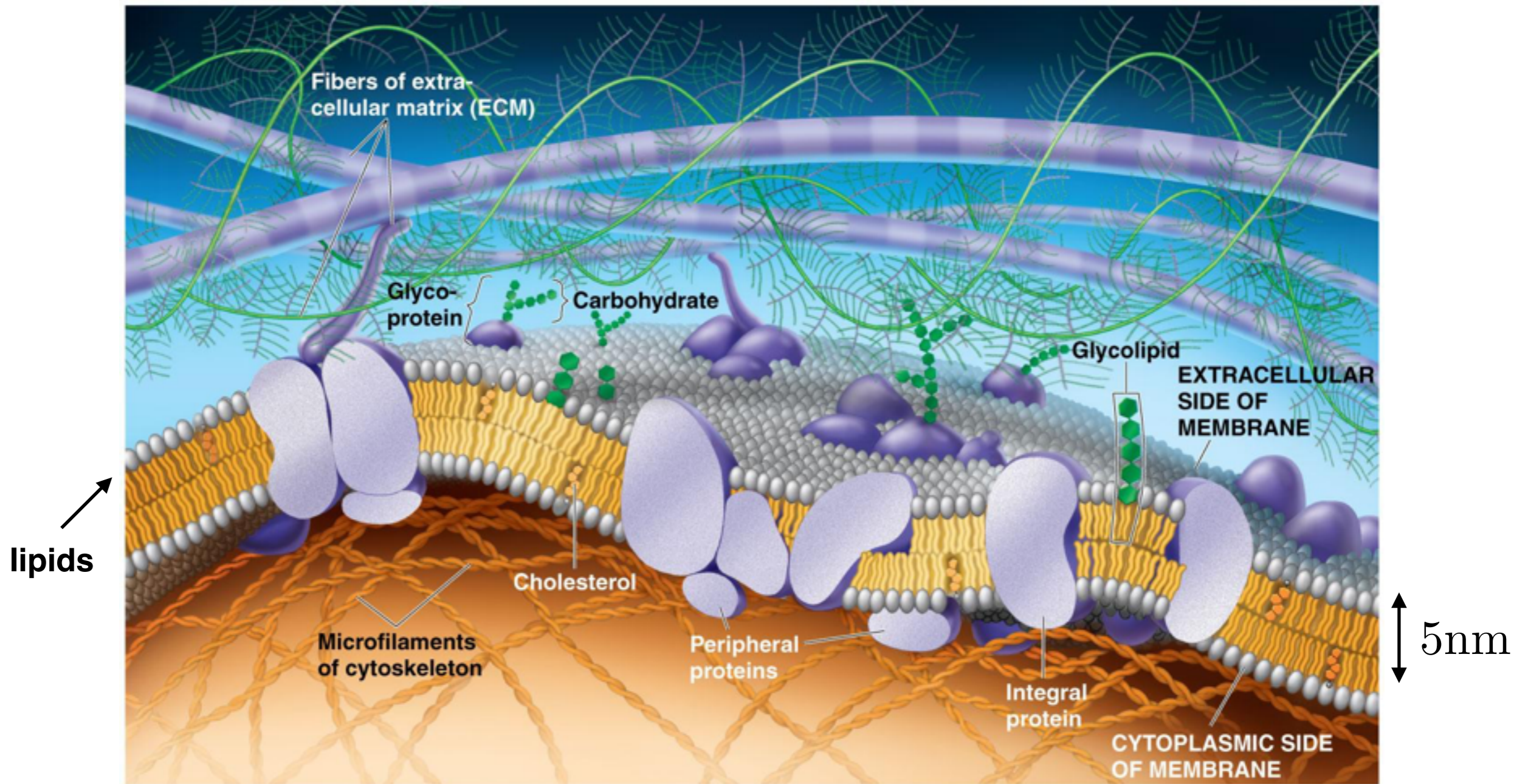


E. Coli



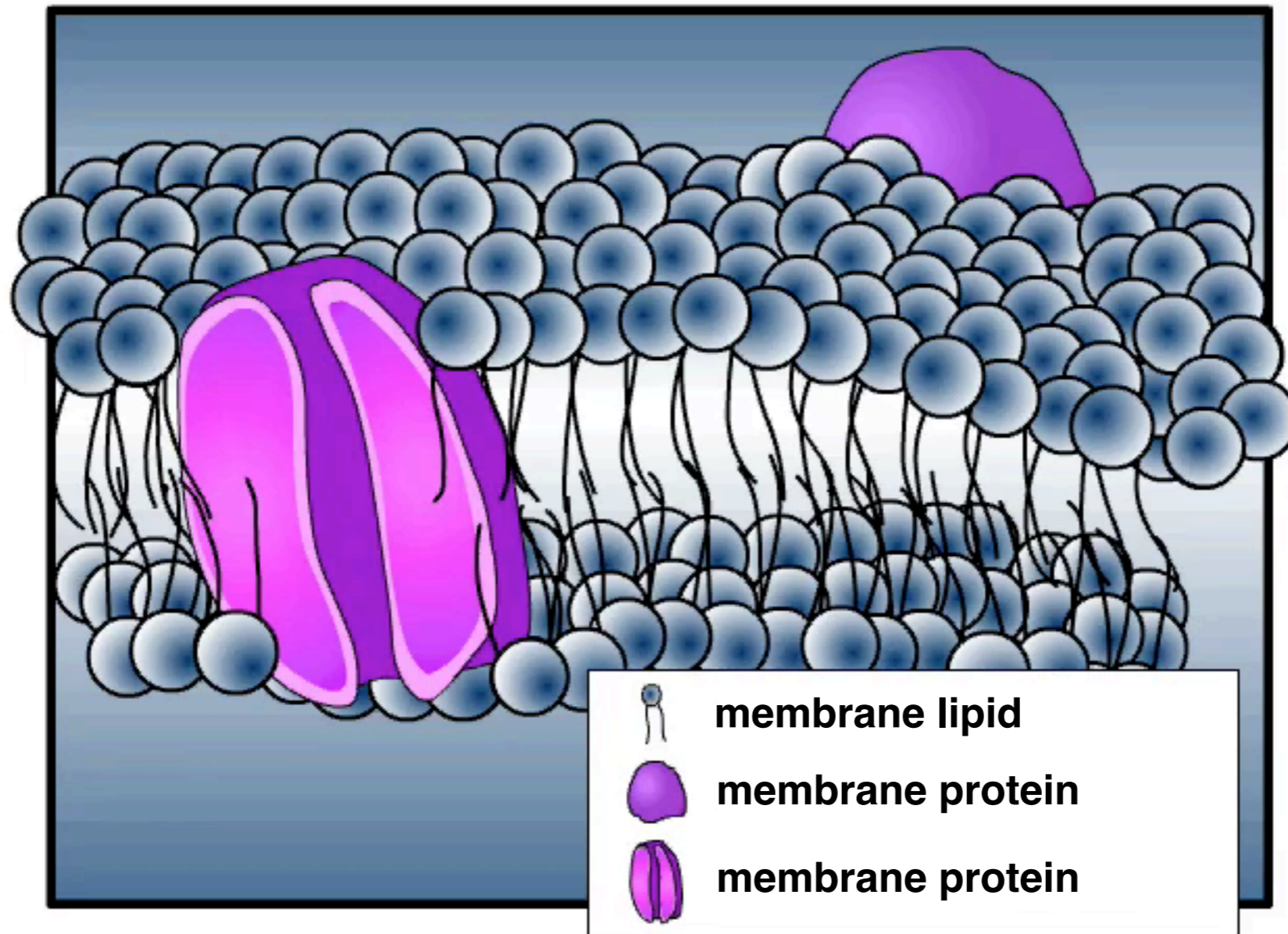
R. Phillips et al., Physical
Biology of the Cell

Cell membrane



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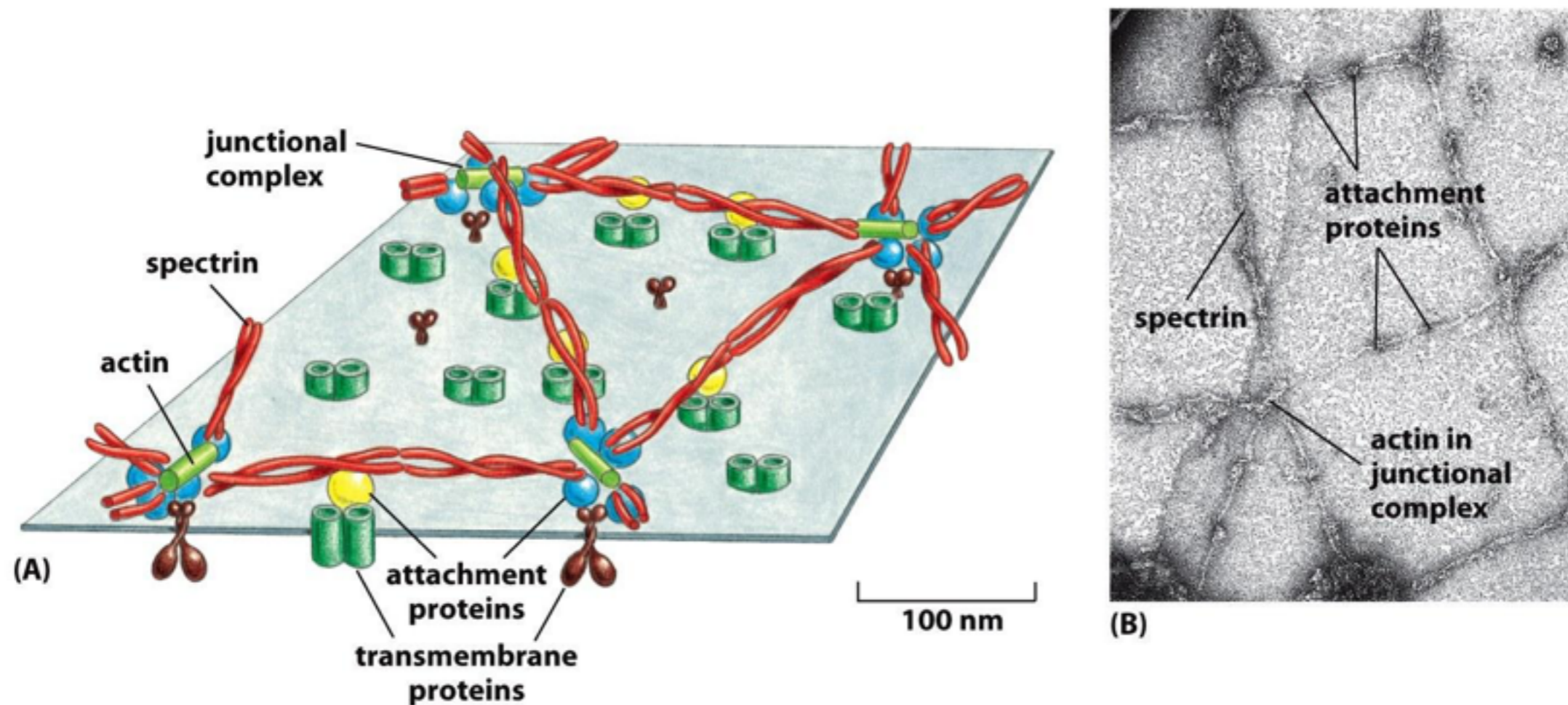
Lipid membrane behaves like fluid



Lipid molecules and proteins can move around!

Flipping of lipid molecules between the layer is unlikely.

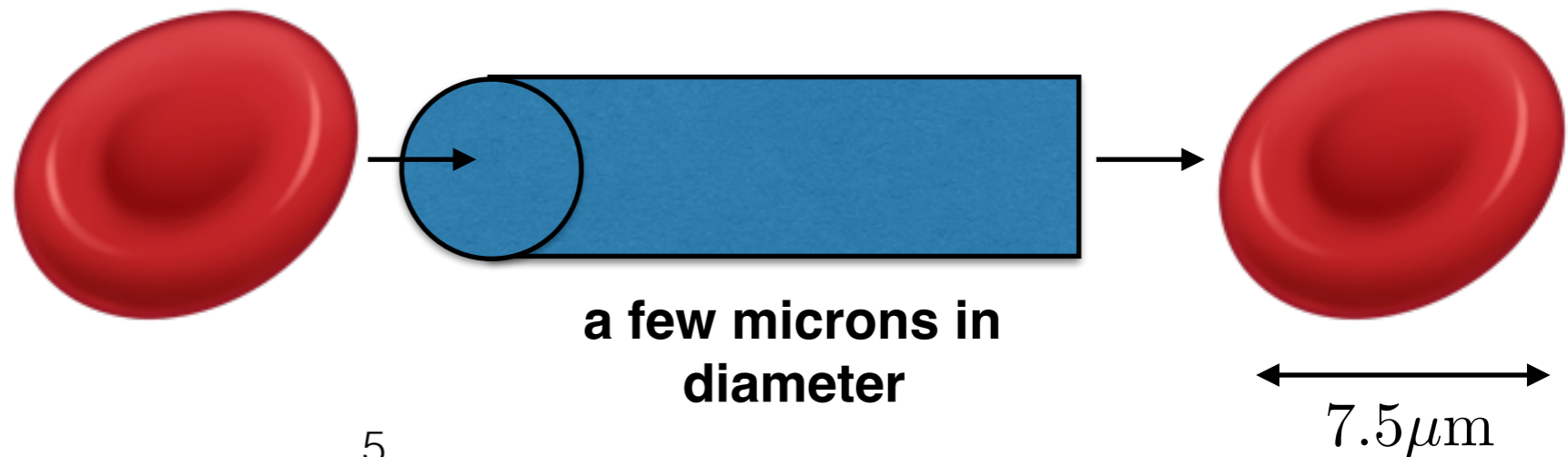
Membrane attached spectrin network provides solid-like behavior



Spectrin network provides structural stability for cells

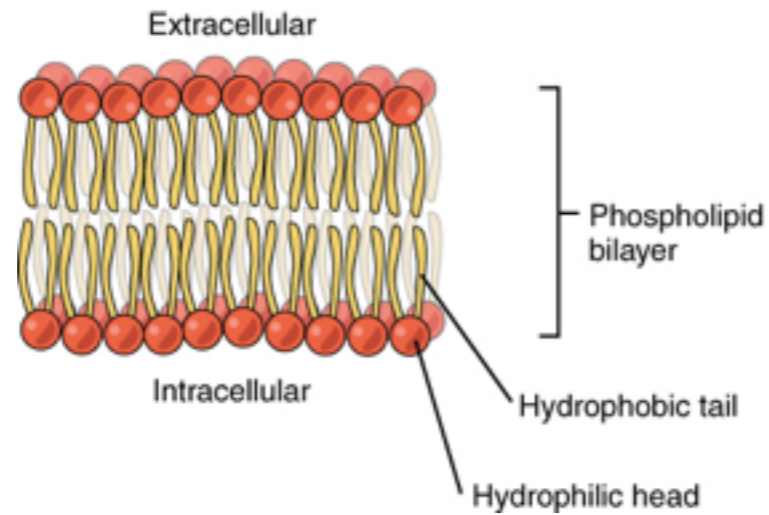
red blood cell

capillary

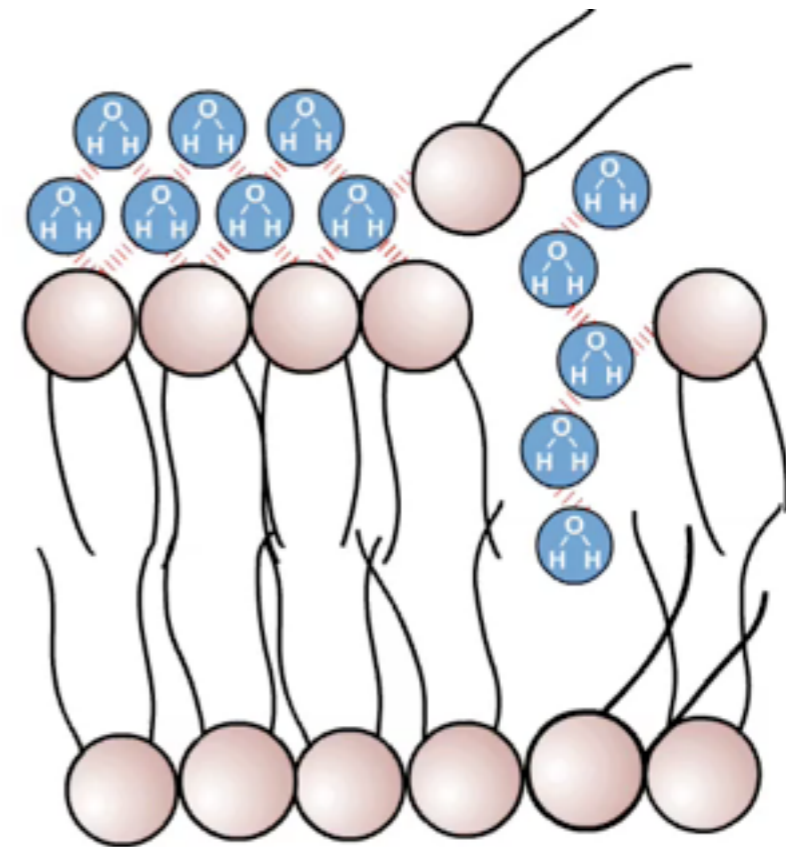
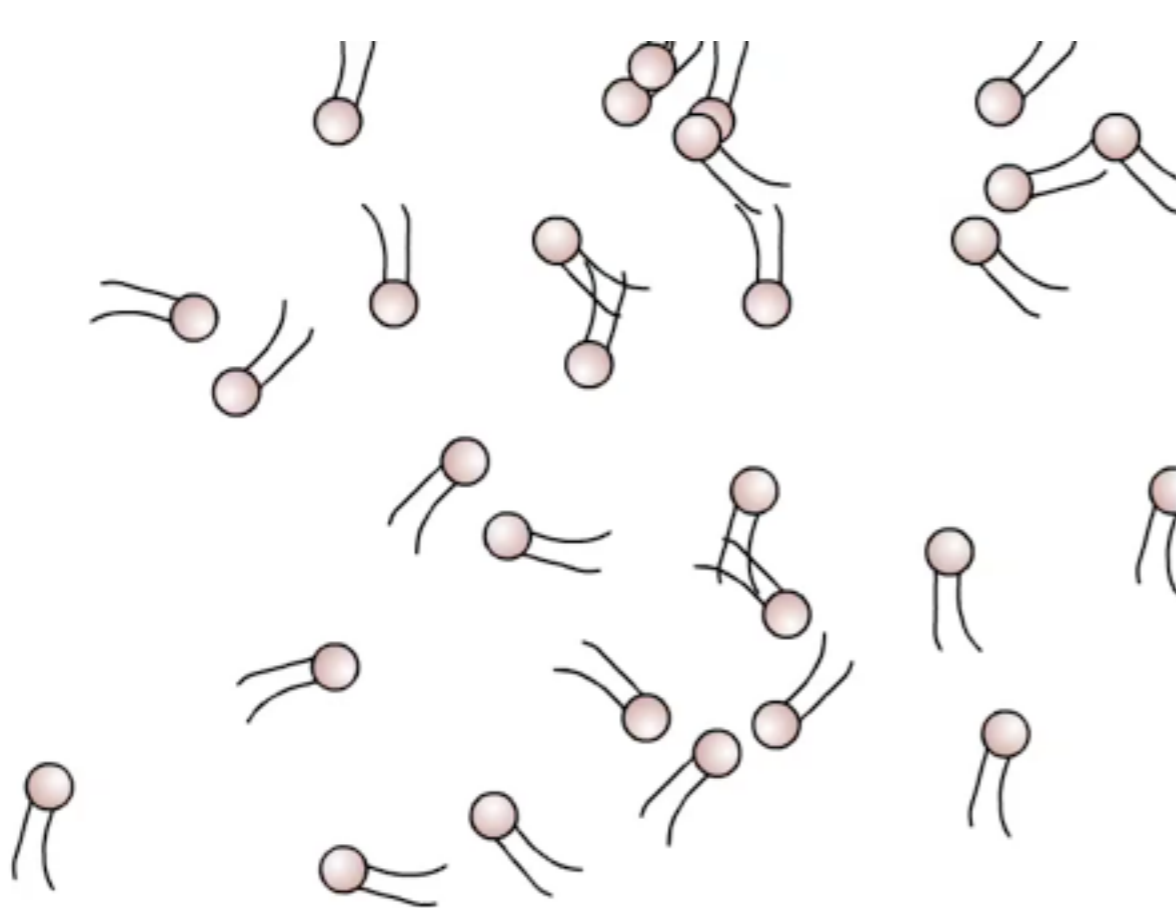


Alberts et al., Molecular Biology of the Cell

Lipid membrane

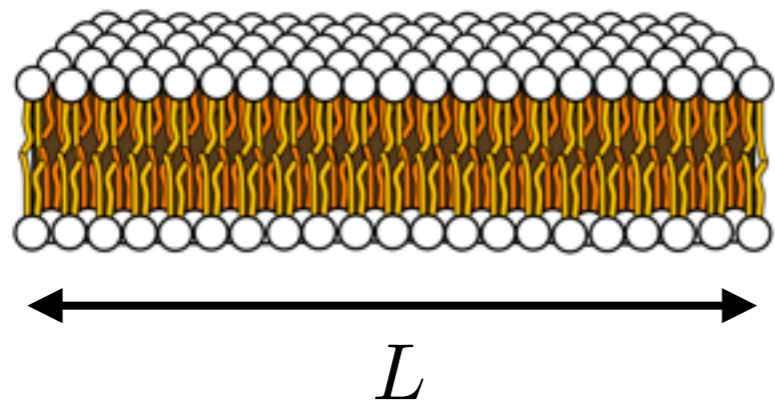


In water solution lipid molecules spontaneously aggregate to prevent undesirable interactions between water and hydrophobic tails.



Flat lipid bilayers vs lipid vesicles

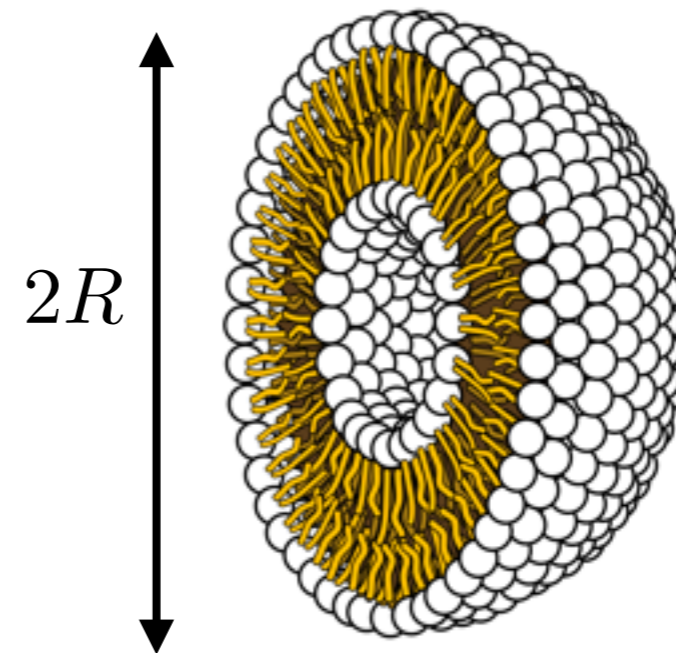
flat bilayer



energy cost on the edge
between lipid tails and
water molecules

$$E \propto L$$

vesicle

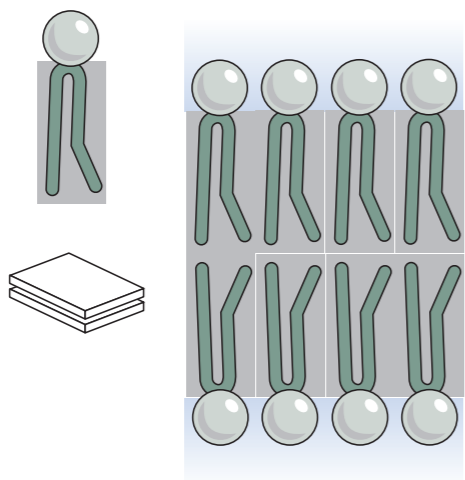


bending energy cost

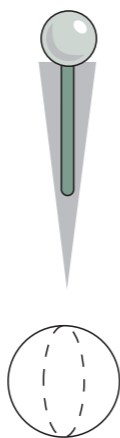
$$E \propto \text{const}$$

Large vesicles have lower energy cost than flat bilayers!

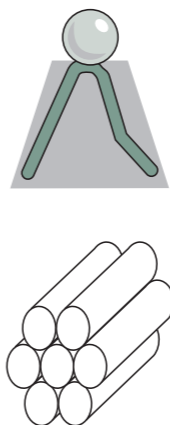
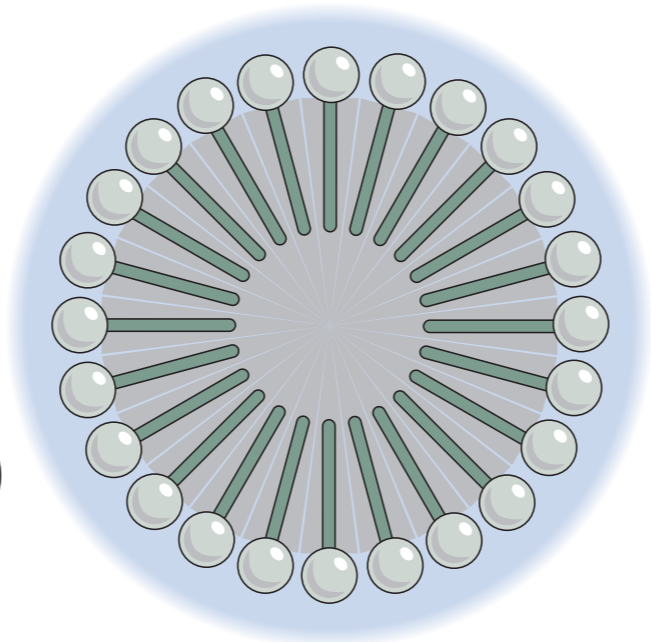
Shape of lipid molecules can induce spontaneous curvature of structures



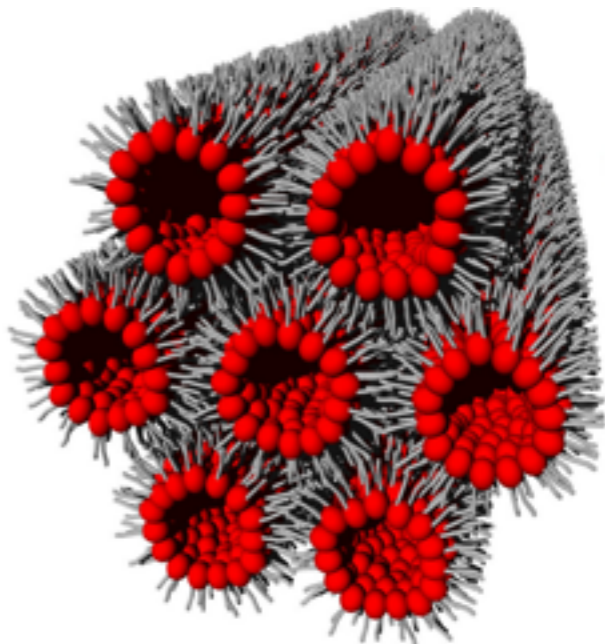
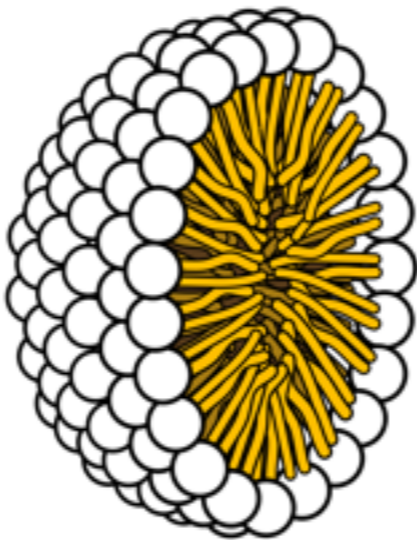
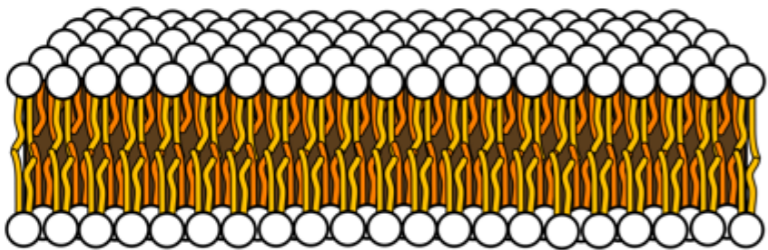
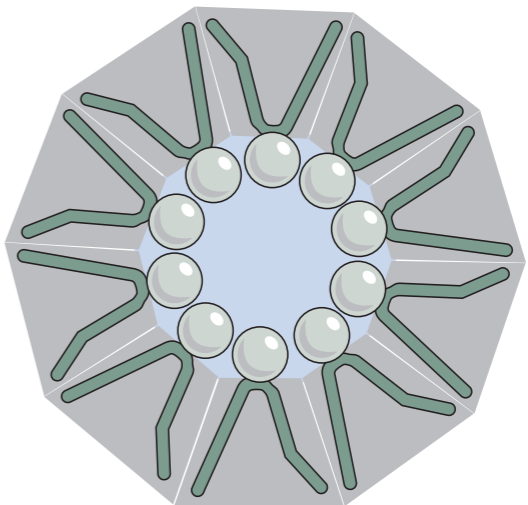
bilayer



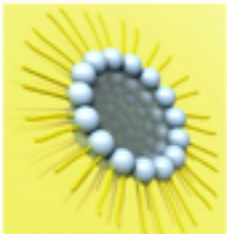
micelle



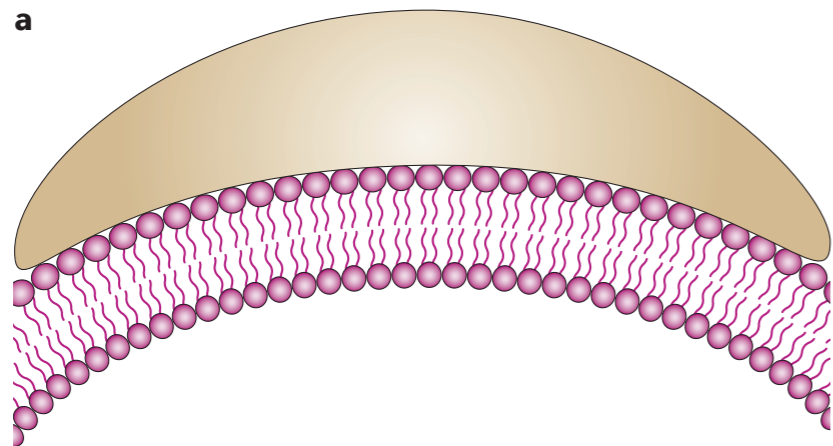
H-II phase



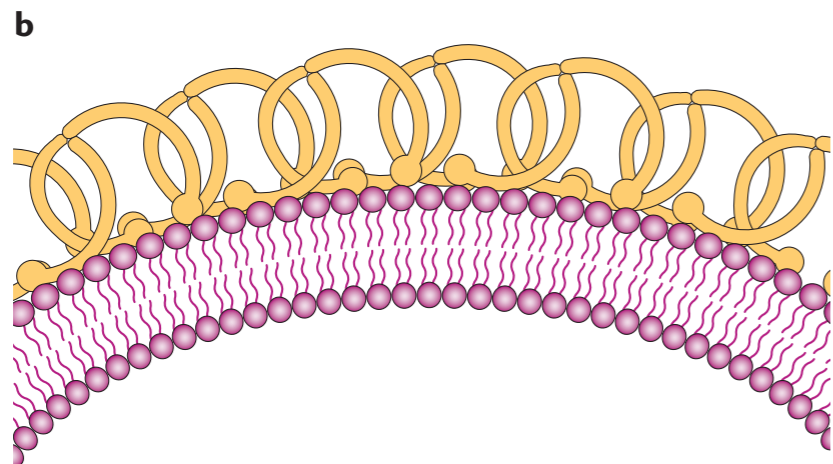
inverted micelle



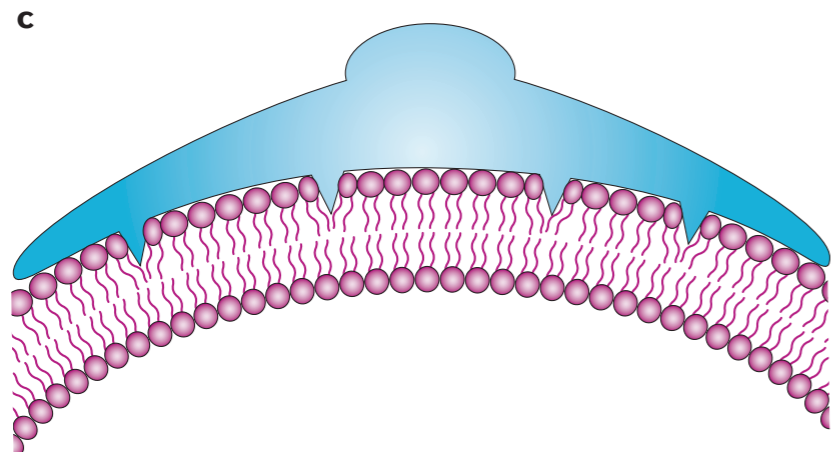
Membrane proteins can induce spontaneous curvature



binding of rigid curved proteins

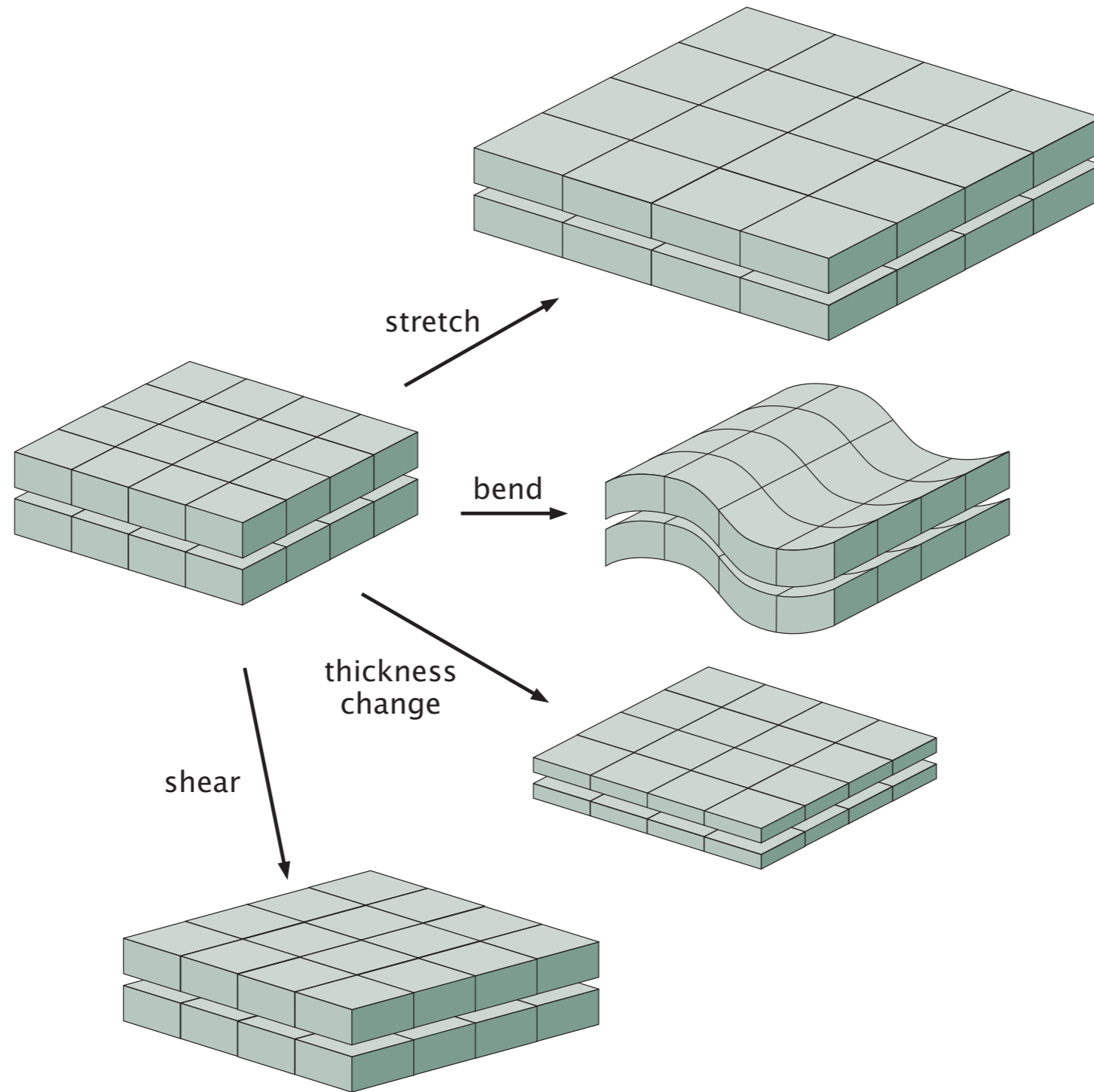


interactions between coat proteins bend the membrane



insertions of protein parts between lipid molecules on one side of the layer

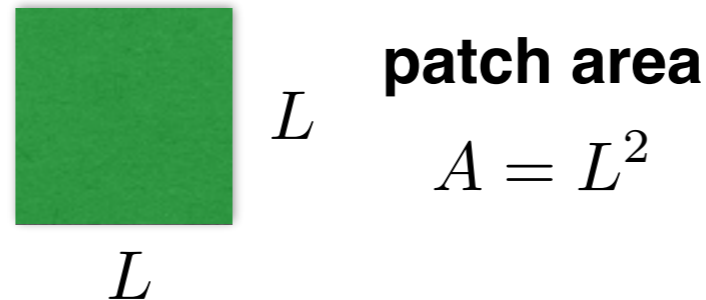
Membrane deformations



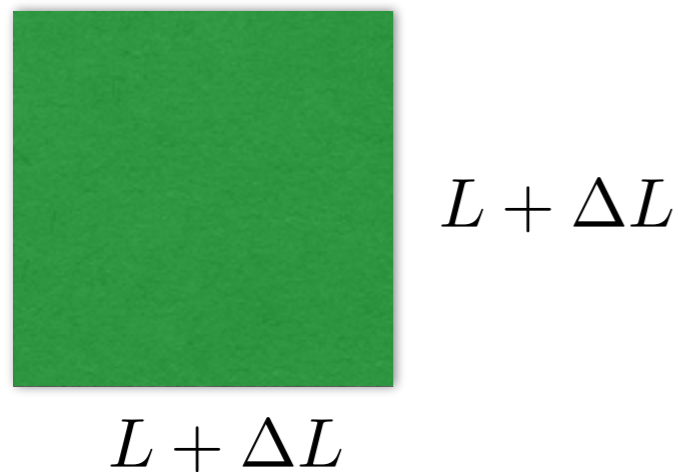
R. Phillips et al., Physical
Biology of the Cell

Energy cost for stretching and shearing

undeformed
square patch



isotropic
deformation



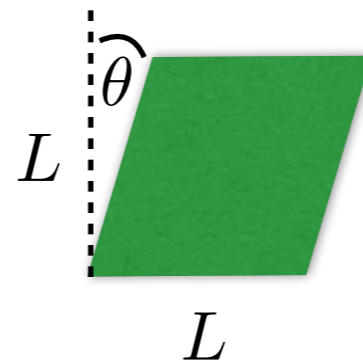
$$\frac{E}{A} = \frac{B}{2} \left(\frac{\Delta A}{A} \right)^2 \approx \frac{B}{2} \left(\frac{2\Delta L}{L} \right)^2$$

bulk modulus

$$B \sim 0.2 \text{ N/m}$$

(lipid bilayer)

shear
deformation



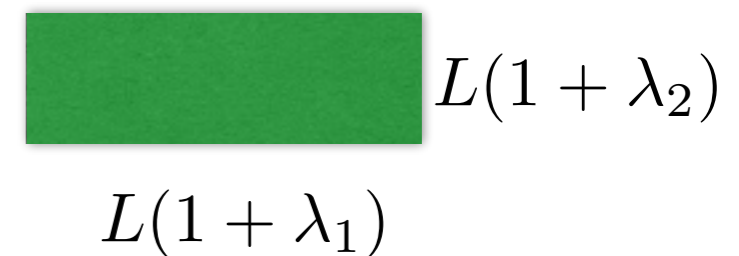
$$\frac{E}{A} = \frac{\mu \theta^2}{2}$$

shear modulus

$$\mu \sim 10^{-5} \text{ N/m}$$

(spectrin network)

anisotropic
stretching



$$\frac{E}{A} \approx \frac{B}{2} (\lambda_1 + \lambda_2)^2 + \frac{\mu}{2} (\lambda_1 - \lambda_2)^2$$

$$\lambda_1, \lambda_2 \ll 1$$

(shearing can be interpreted
as anisotropic stretching)